

ADA Homework Assignment #1

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Challenge 1

The R code below will create a vector called **quote** that contains the last line of Charles Darwin's *On the Origin of Species*

```
quote <- ("There is grandeur in this view of life, with its several powers, having been originally breathe  
quote
```

```
## [1] "There is grandeur in this view of life, with its several powers, having been originally breathe
```

The following will split the string of characters in the vector called **quote** into individual words while also removing any punctuation contained in the vector. Loading the {stringr} package is necessary here.

```
library(stringr)  
split_quote <- str_split((gsub("[:punct:]", "", quote)), " ")  
split_quote
```

```
## [[1]]  
## [1] "There"      "is"         "grandeur"   "in"         "this"  
## [6] "view"       "of"         "life"       "with"       "its"  
## [11] "several"    "powers"     "having"     "been"       "originally"  
## [16] "breathed"   "by"         "the"        "Creator"    "into"  
## [21] "a"          "few"        "forms"      "or"         "into"  
## [26] "one"        "and"        "that"       "whilst"     "this"  
## [31] "planet"     "has"        "gone"       "circling"   "on"  
## [36] "according"  "to"         "the"        "fixed"      "law"  
## [41] "of"         "gravity"    "from"       "so"         "simple"  
## [46] "a"          "beginning" "endless"    "forms"      "most"  
## [51] "beautiful" "and"        "most"       "wonderful"  "have"  
## [56] "been"       "and"        "are"        "being"      "evolved"
```

The following R code creates a new vector of every fourth entry from the vector contained in the list of the quote split into individual words

```
every_fourth<-split_quote[[1]][1:15*4]  
every_fourth
```

```
## [1] "in"         "life"       "powers"     "breathed"   "into"       "or"  
## [7] "that"       "has"        "according"  "law"        "so"         "endless"  
## [13] "and"        "been"       "evolved"
```

The below code will sort the entries in the defined list in reverse alphabetical order

```
sort(every_fourth, decreasing=TRUE)
```

```
## [1] "that"      "so"      "powers"   "or"      "life"     "law"
## [7] "into"      "in"      "has"      "evolved" "endless"  "breathed"
## [13] "been"     "and"     "according"
```

Challenge 2

Creation of an 8 x 10 matrix called **m1** consisting of odd integers from 1 to 159 filled column-wise

```
m1 <- matrix(data=seq(from=1,to=159,by=2), nrow=8, ncol=10)
m1
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
## [1,]    1   17   33   49   65   81   97  113  129  145
## [2,]    3   19   35   51   67   83   99  115  131  147
## [3,]    5   21   37   53   69   85  101  117  133  149
## [4,]    7   23   39   55   71   87  103  119  135  151
## [5,]    9   25   41   57   73   89  105  121  137  153
## [6,]   11   27   43   59   75   91  107  123  139  155
## [7,]   13   29   45   61   77   93  109  125  141  157
## [8,]   15   31   47   63   79   95  111  127  143  159
```

Extraction of the element in row 5, column 2 using single-bracket notation

```
m1[5,2]
```

```
## [1] 25
```

Extraction of rows 5 to 7 in the matrix **m1**

```
m1[5:7,]
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
## [1,]    9   25   41   57   73   89  105  121  137  153
## [2,]   11   27   43   59   75   91  107  123  139  155
## [3,]   13   29   45   61   77   93  109  125  141  157
```

Creation of a new variable **m2** that contains (row 3, column 4) to (row 6, column 9) of **m1** as well as the code to describe the class and mode of **m2**

```
m1$m2 <- m1[3:6, 4:9]
```

```
## Warning in m1$m2 <- m1[3:6, 4:9]: Coercing LHS to a list
```

```
m1$m2
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6]
## [1,]   53   69   85  101  117  133
## [2,]   55   71   87  103  119  135
## [3,]   57   73   89  105  121  137
## [4,]   59   75   91  107  123  139
```

```
class(m1$m2)
```

```
## [1] "matrix"
```

```
mode(m1$m2)
```

```
## [1] "numeric"
```

Challenge 3

Constructs a 4-dimensional, 400-element array (5 x 5 x 4 x 4), **a**, consisting of the numbers 400 to 1

```
a <- array(data=400:1, dim= c(5,5,4,4))
a
```

```
## , , 1, 1
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,]  400  395  390  385  380
## [2,]  399  394  389  384  379
## [3,]  398  393  388  383  378
## [4,]  397  392  387  382  377
## [5,]  396  391  386  381  376
##
## , , 2, 1
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,]  375  370  365  360  355
## [2,]  374  369  364  359  354
## [3,]  373  368  363  358  353
## [4,]  372  367  362  357  352
## [5,]  371  366  361  356  351
##
## , , 3, 1
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,]  350  345  340  335  330
## [2,]  349  344  339  334  329
## [3,]  348  343  338  333  328
## [4,]  347  342  337  332  327
## [5,]  346  341  336  331  326
##
```

```

## , , 4, 1
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 325 320 315 310 305
## [2,] 324 319 314 309 304
## [3,] 323 318 313 308 303
## [4,] 322 317 312 307 302
## [5,] 321 316 311 306 301
##
## , , 1, 2
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 300 295 290 285 280
## [2,] 299 294 289 284 279
## [3,] 298 293 288 283 278
## [4,] 297 292 287 282 277
## [5,] 296 291 286 281 276
##
## , , 2, 2
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 275 270 265 260 255
## [2,] 274 269 264 259 254
## [3,] 273 268 263 258 253
## [4,] 272 267 262 257 252
## [5,] 271 266 261 256 251
##
## , , 3, 2
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 250 245 240 235 230
## [2,] 249 244 239 234 229
## [3,] 248 243 238 233 228
## [4,] 247 242 237 232 227
## [5,] 246 241 236 231 226
##
## , , 4, 2
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 225 220 215 210 205
## [2,] 224 219 214 209 204
## [3,] 223 218 213 208 203
## [4,] 222 217 212 207 202
## [5,] 221 216 211 206 201
##
## , , 1, 3
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 200 195 190 185 180
## [2,] 199 194 189 184 179
## [3,] 198 193 188 183 178
## [4,] 197 192 187 182 177
## [5,] 196 191 186 181 176
##

```

```

## , , 2, 3
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 175 170 165 160 155
## [2,] 174 169 164 159 154
## [3,] 173 168 163 158 153
## [4,] 172 167 162 157 152
## [5,] 171 166 161 156 151
##
## , , 3, 3
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 150 145 140 135 130
## [2,] 149 144 139 134 129
## [3,] 148 143 138 133 128
## [4,] 147 142 137 132 127
## [5,] 146 141 136 131 126
##
## , , 4, 3
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 125 120 115 110 105
## [2,] 124 119 114 109 104
## [3,] 123 118 113 108 103
## [4,] 122 117 112 107 102
## [5,] 121 116 111 106 101
##
## , , 1, 4
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 100 95 90 85 80
## [2,] 99 94 89 84 79
## [3,] 98 93 88 83 78
## [4,] 97 92 87 82 77
## [5,] 96 91 86 81 76
##
## , , 2, 4
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 75 70 65 60 55
## [2,] 74 69 64 59 54
## [3,] 73 68 63 58 53
## [4,] 72 67 62 57 52
## [5,] 71 66 61 56 51
##
## , , 3, 4
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,] 50 45 40 35 30
## [2,] 49 44 39 34 29
## [3,] 48 43 38 33 28
## [4,] 47 42 37 32 27
## [5,] 46 41 36 31 26
##

```

```
## , , 4, 4
##
##      [,1] [,2] [,3] [,4] [,5]
## [1,]   25   20   15   10    5
## [2,]   24   19   14    9    4
## [3,]   23   18   13    8    3
## [4,]   22   17   12    7    2
## [5,]   21   16   11    6    1
```

Extractions of varying elements of parts of **a**

```
a[1,1,1,2]
```

```
## [1] 300
```

```
a[2,3,2,]
```

```
## [1] 364 264 164 64
```

```
a[1:5,1:5,3,3]
```

```
##      [,1] [,2] [,3] [,4] [,5]
## [1,]  150  145  140  135  130
## [2,]  149  144  139  134  129
## [3,]  148  143  138  133  128
## [4,]  147  142  137  132  127
## [5,]  146  141  136  131  126
```

Challenge 4

The following creates a list of the (simplified) primate taxonomy including the taxonomic levels and names of each internal node.

```
Lorisoidea <- list("Lorisidae", "Galagidae")
Lorisiformes <- list(Lorisoidea)
names(Lorisiformes)<- c("Superfamily_Lorisoidea")
Lemuroidea <- list("Cheirogaleidae", "Lepilemuridae", "Indriidae", "Lemuridae", "Daubentoniidae")
Lemuriformes <- list(Lemuroidea)
names(Lemuriformes)<- c("Superfamily_Lemuroidea")
Strepsirhini <- list(Lemuriformes, Lorisiformes)
names(Strepsirhini) <- c("Infraorder_Lemuriformes", "Infraorder_Lorisiformes")
Cercopithecoidea <- list("Cercopithecidae")
Hominoidea <- list("Hylobatidae", "Hominidae")
Catarrhini <- list(Hominoidea, Cercopithecoidea)
names(Catarrhini)<- c("Superfamily_Hominoidea", "Superfamily_Cercopithecoidea")
Ceboidea <- list("Cebidae", "Atelidae", "Pitheciidae")
Platyrrhini <- list(Ceboidea)
names(Platyrrhini)<-c("Superfamily_Ceboidea")
Simiiformes <- list(Platyrrhini, Catarrhini)
names(Simiiformes)<-c("Parvorder_Platyrrhini", "Parvorder_Catarrhini")
Tarsioidea <- list("Tarsiidae")
```

```
Tarsiiformes <- list(Tarsiodea)
names(Tarsiiformes)<-c("Superfamily_Tarsiodea")
Haplorhini <- list(Tarsiiformes, Simiiformes)
names(Haplorhini)<-c("Infraorder_Tarsiiformes","Infraorder_Simiiformes")
Primates <- list(Strepsirhini, Haplorhini)
names(Primates)<-c("Suborder_Strepsirhini","Suborder_Haplorhini")
```

The following uses double bracket notation to extract the New World Monkeys (Parvorder: Platyrrhini) from the list as a new variable **nwm** and lists the class and mode of the variable.

```
Primates$nwm <- Primates[[2]][[2]][1]
Primates$nwm
```

```
## $Parvorder_Platyrrhini
## $Parvorder_Platyrrhini$Superfamily_Ceboidea
## $Parvorder_Platyrrhini$Superfamily_Ceboidea[[1]]
## [1] "Cebidae"
##
## $Parvorder_Platyrrhini$Superfamily_Ceboidea[[2]]
## [1] "Atelidae"
##
## $Parvorder_Platyrrhini$Superfamily_Ceboidea[[3]]
## [1] "Pitheciidae"
```

```
class(Primates$nwm)
```

```
## [1] "list"
```

```
mode(Primates$nwm)
```

```
## [1] "list"
```

The following uses \$ notation to extract the tarsiers from the list

```
Primates$Suborder_Haplorhini$Infraorder_Tarsiiformes
```

```
## $Superfamily_Tarsiodea
## $Superfamily_Tarsiodea[[1]]
## [1] "Tarsiidae"
```